\_\_\_\_\_\_

Sequence Listing was accepted with existing errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: Thu May 17 08:56:14 EDT 2007

\_\_\_\_\_

## Validated By CRFValidator v 1.0.2

Application No: 10564088 Version No: 1.1

Input Set:

Output Set:

**Started:** 2007-05-17 08:56:03.380

**Finished:** 2007-05-17 08:56:03.679

**Elapsed:** 0 hr(s) 0 min(s) 0 sec(s) 299 ms

Total Warnings: 4

Total Errors: 0

No. of SeqIDs Defined: 6

Actual SeqID Count: 6

| Error code |     | Error Description |    |         |       |    |       |    |     |    |     |  |  |  |
|------------|-----|-------------------|----|---------|-------|----|-------|----|-----|----|-----|--|--|--|
| W          | 213 | Artificial        | or | Unknown | found | in | <213> | in | SEQ | ID | (3) |  |  |  |
| W          | 213 | Artificial        | or | Unknown | found | in | <213> | in | SEQ | ID | (4) |  |  |  |
| W          | 213 | Artificial        | or | Unknown | found | in | <213> | in | SEQ | ID | (5) |  |  |  |
| W          | 213 | Artificial        | or | Unknown | found | in | <213> | in | SEQ | ID | (6) |  |  |  |

## SEQUENCE LISTING <110> ZGene A/S Gojkovic, Zoran <120> Yellow fever mosquito deoxyribonucleoside kinase and its use <130> 519-204-WO <140> 10/564,088 <141> 2006-01-11 <150> DK PA 2003 01067 <151> 2003-07-11 <160> 6 <170> PatentIn version 3.1 <210> 1 <211> 747 <212> DNA <213> Aedes aegypti <220>

| <220>   |               |     |     |     |     |     |     |     |     |     |     |     |     |     |  |
|---------|---------------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| <221>   | > CDS         |     |     |     |     |     |     |     |     |     |     |     |     |     |  |
| <222>   | 222> (1)(747) |     |     |     |     |     |     |     |     |     |     |     |     |     |  |
| <400> 1 |               |     |     |     |     |     |     |     |     |     |     |     |     |     |  |
| atg gcg | gct           | gcc | atc | gga | ccg | gag | cgg | ctt | ggc | gtg | gcc | gga | aag | aag |  |
| Met Ala | Ala           | Ala | Ile | Gly | Pro | Glu | Arg | Leu | Gly | Val | Ala | Gly | Lys | Lys |  |
| 1       |               |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |  |
|         |               |     |     |     |     |     |     |     |     |     |     |     |     |     |  |
| ccc ttc | act           | gtt | ttc | att | gag | gga | aac | atc | ggc | agc | gga | aag | acc | aca |  |

20

|     | _   |     |     |     |     |     |     | _   | -   |     | _   | _   | _   | _   | _   |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Phe | Leu | Asn | His | Phe | Glu | Lys | Phe | Lys | Asp | Arg | Val | Cys | Leu | Leu | Thr |     |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| gaa | cct | gtg | gaa | aag | tgg | cgg | gac | tgc | ggg | gga | gtc | aat | ctg | ctg | gat | 192 |
| Glu | Pro | Val | Glu | Lys | Trp | Arg | Asp | Cys | Gly | Gly | Val | Asn | Leu | Leu | Asp |     |
|     | 50  |     |     |     | _   | 55  |     |     |     |     | 60  |     |     |     | _   |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| cta | atg | tac | aag | gaa | ccg | cac | cgt | tgg | gcg | atg | ccg | ttc | cag | acc | tac | 240 |

Pro Phe Thr Val Phe Ile Glu Gly Asn Ile Gly Ser Gly Lys Thr Thr

ttc ctg aac cac ttc gag aaa ttc aag gat agg gtt tgt ctg ctg acg

48

96

144

cta atg tac aag gaa ccg cac cgt tgg gcg atg ccg ttc cag acc tac Leu Met Tyr Lys Glu Pro His Arg Trp Ala Met Pro Phe Gln Thr Tyr 65 70 75 80

| _   |     |     | _          | atg<br>Met        |     |     | _   |            | Thr       |     |     | _   | -          | Lys       | _   | 288 |
|-----|-----|-----|------------|-------------------|-----|-----|-----|------------|-----------|-----|-----|-----|------------|-----------|-----|-----|
| gtg | aag | ctc | atg        | 85<br>gaa         | cgg | tcc | atg | ttc        | 90<br>agt | gcc | aga | tat | tgt        | 95<br>ttc | gtg | 336 |
| Val | Lys | Leu | Met<br>100 | Glu               | Arg | Ser | Met | Phe<br>105 | Ser       | Ala | Arg | Tyr | Cys<br>110 | Phe       | Val |     |
|     |     | Met |            | gcg<br>Ala        |     |     | Ser |            |           |     |     | Met |            |           |     | 384 |
| cta | caa | 115 | tgg        | tat               | gag | ttc | 120 | cat        | qcc       | aat | ata | 125 | att        | caa       | gca | 432 |
|     |     |     |            | Tyr               |     |     |     |            | -         |     |     |     |            |           | _   |     |
| _   |     |     | -          | tat<br>Tyr        |     | _   |     | _          | _         | _   |     | _   |            |           | _   | 480 |
| 145 |     |     |            |                   | 150 |     |     |            |           | 155 |     |     |            |           | 160 | 500 |
| _   |     | _   | -          | gca<br>Ala<br>165 | -   | -   | -   | -          | -         | _   | -   | _   |            |           |     | 528 |
|     |     | -   |            | cac<br>His        |     | _   |     | -          |           |     |     |     |            |           |     | 576 |
|     | _   | _   | _          | gcc<br>Ala        | _   | _   |     | _          | _         | _   | _   | _   |            | -         | _   | 624 |
|     |     |     | -          | tca<br>Ser        | -   |     | -   | -          |           | -   |     | -   |            |           | -   | 672 |
|     |     |     |            | gat<br>Asp        |     |     |     | _          |           |     |     |     | -          |           |     | 720 |
| _   |     | _   | _          | aga<br>Arg<br>245 |     | _   |     | taa        |           |     |     |     |            |           |     | 747 |

<210> 2

<211> 248

<212> PRT

<213> Aedes aegypti

Met Ala Ala Ala Ile Gly Pro Glu Arg Leu Gly Val Ala Gly Lys Lys Pro Phe Thr Val Phe Ile Glu Gly Asn Ile Gly Ser Gly Lys Thr Thr 20 25 Phe Leu Asn His Phe Glu Lys Phe Lys Asp Arg Val Cys Leu Leu Thr Glu Pro Val Glu Lys Trp Arg Asp Cys Gly Gly Val Asn Leu Leu Asp 55 Leu Met Tyr Lys Glu Pro His Arg Trp Ala Met Pro Phe Gln Thr Tyr 70 75 Val Thr Leu Thr Met Leu Asn Met His Thr Tyr Gln Thr Asp Lys Ser 85 90 Val Lys Leu Met Glu Arg Ser Met Phe Ser Ala Arg Tyr Cys Phe Val 105 100 Glu Asn Met Leu Ala Ser Gly Ser Leu His Gln Gly Met Tyr Asn Ile 115 120 Leu Gln Glu Trp Tyr Glu Phe Ile His Ala Asn Ile His Ile Gln Ala 135 140 Asp Leu Ile Val Tyr Leu Arg Thr Ser Pro Glu Ile Val Tyr Glu Arg 150 155 Met Lys Lys Arg Ala Arg Ser Glu Glu Ser Cys Val Pro Leu Lys Tyr 165 170 Leu Gln Glu Leu His Glu Leu His Glu Asn Trp Leu Ile His Gly Thr 180 Phe Pro Arg Val Ala Pro Val Leu Val Leu Asp Ala Asp Leu Asp Leu 195 200 His Asn Ile Ser Ser Glu Tyr Lys Arg Ser Glu Thr Ser Ile Leu Lys 210 215 220 Pro Ile Leu Ile Asp Asn Thr Asn Gln His Pro Ile Leu Ala Ser Pro 230 235 Ser Lys Arg Ser Arg Thr Glu Phe 245 <210> 3

<211> 28

<212> DNA

<213> Artificial sequence

```
<220>
<223> PCR primer sequence
<400> 3
ttaggatcca tggcggctgc catcggac
                                                                    28
<210> 4
<211> 30
<212> DNA
<213> artificial sequence
<220>
<223> PCR primer sequence
<400> 4
                                                                     30
cagcaattgt tagaattcag ttctcgatcg
<210> 5
<211> 31
<212> DNA
<213> artificial sequence
<220>
<223> PCR primer sequence
<400> 5
cgcggatcca tggcttcgta ccccggccat c
                                                                    31
<210> 6
<211> 33
<212> DNA
<213> artificial sequence
```

<223> PCR primer sequence

<400> 6 ccggaattct tagttagcct ccccatctc ccg

33